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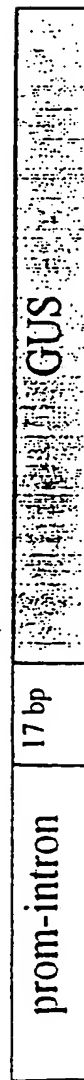
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PROMOTER REGION	REPORTER GENE	CONSTRUCTS
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86AGUS



85-17 GUS



Δ GUS

FIG. 1

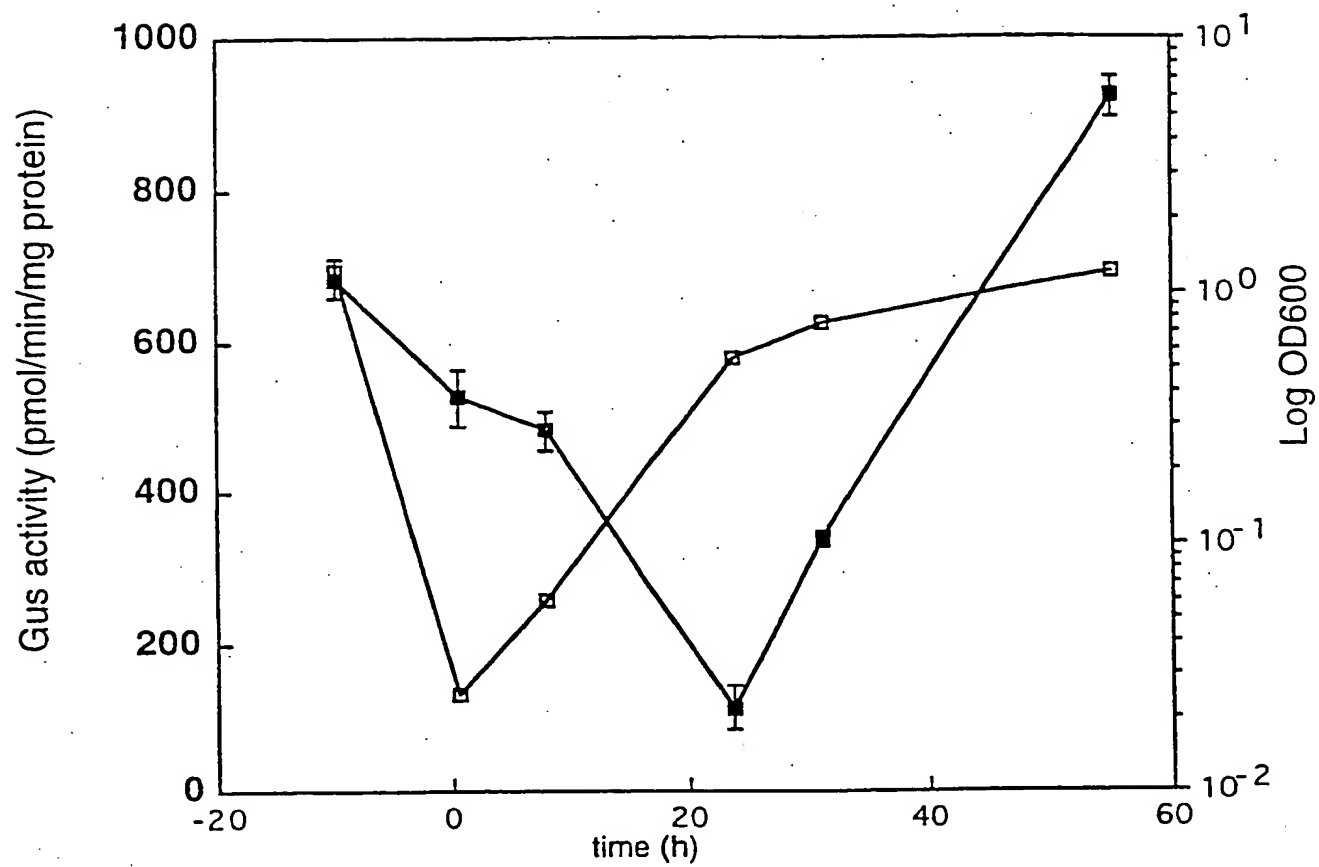
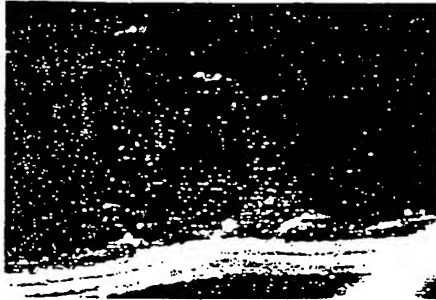
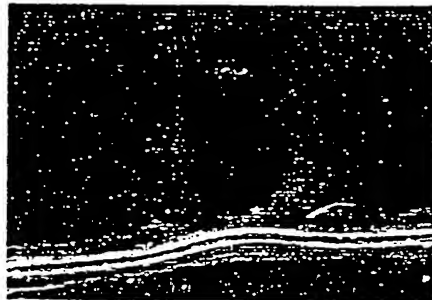


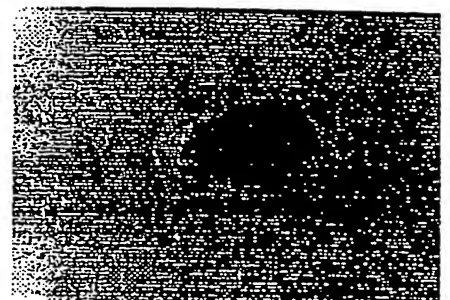
FIG. 2



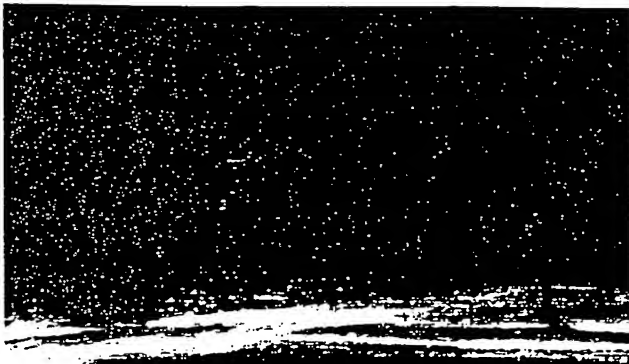
A



B



C



D



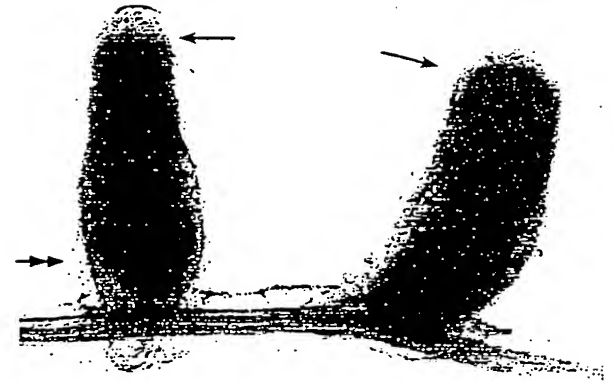
E



F



G



H

FIG. 3

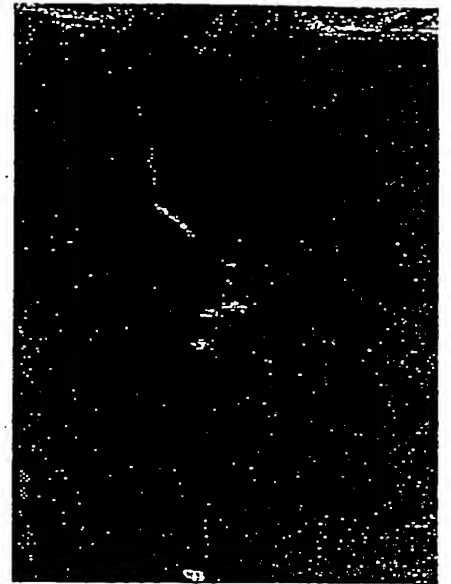
FIG. 4



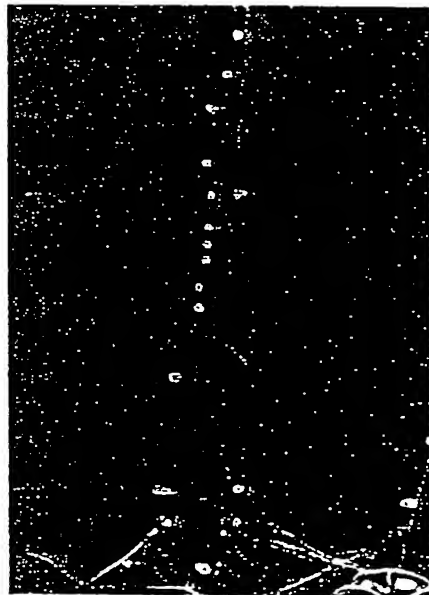
A



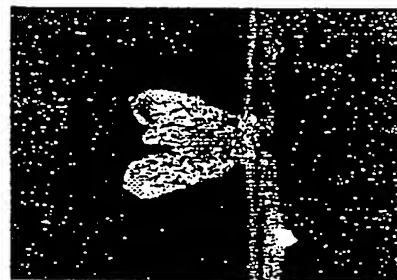
B



C



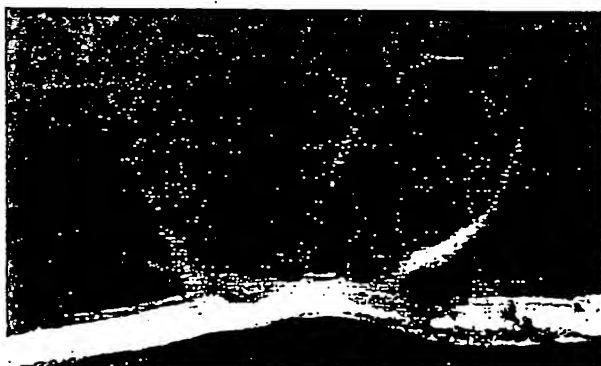
D



E1



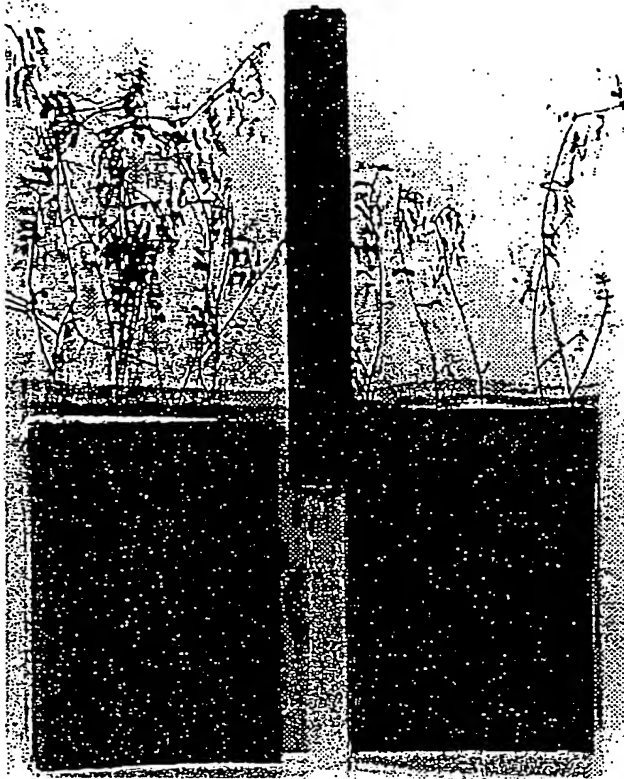
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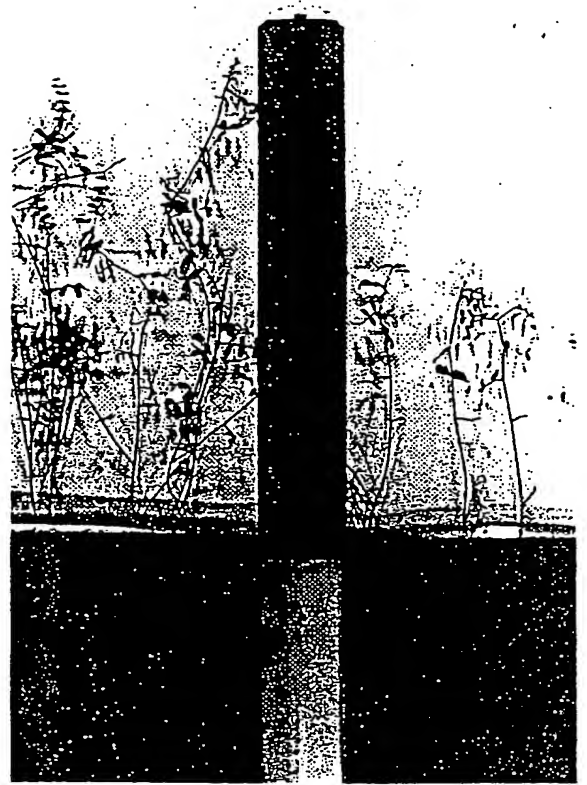
F



G



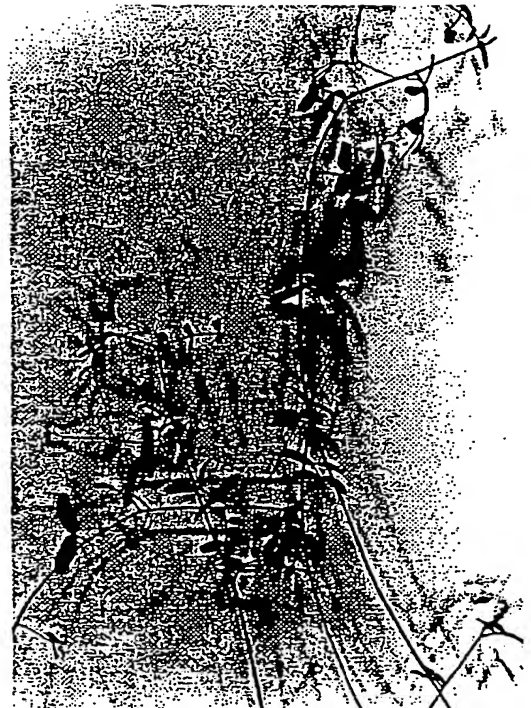
A



B



C



D

FIG. 5

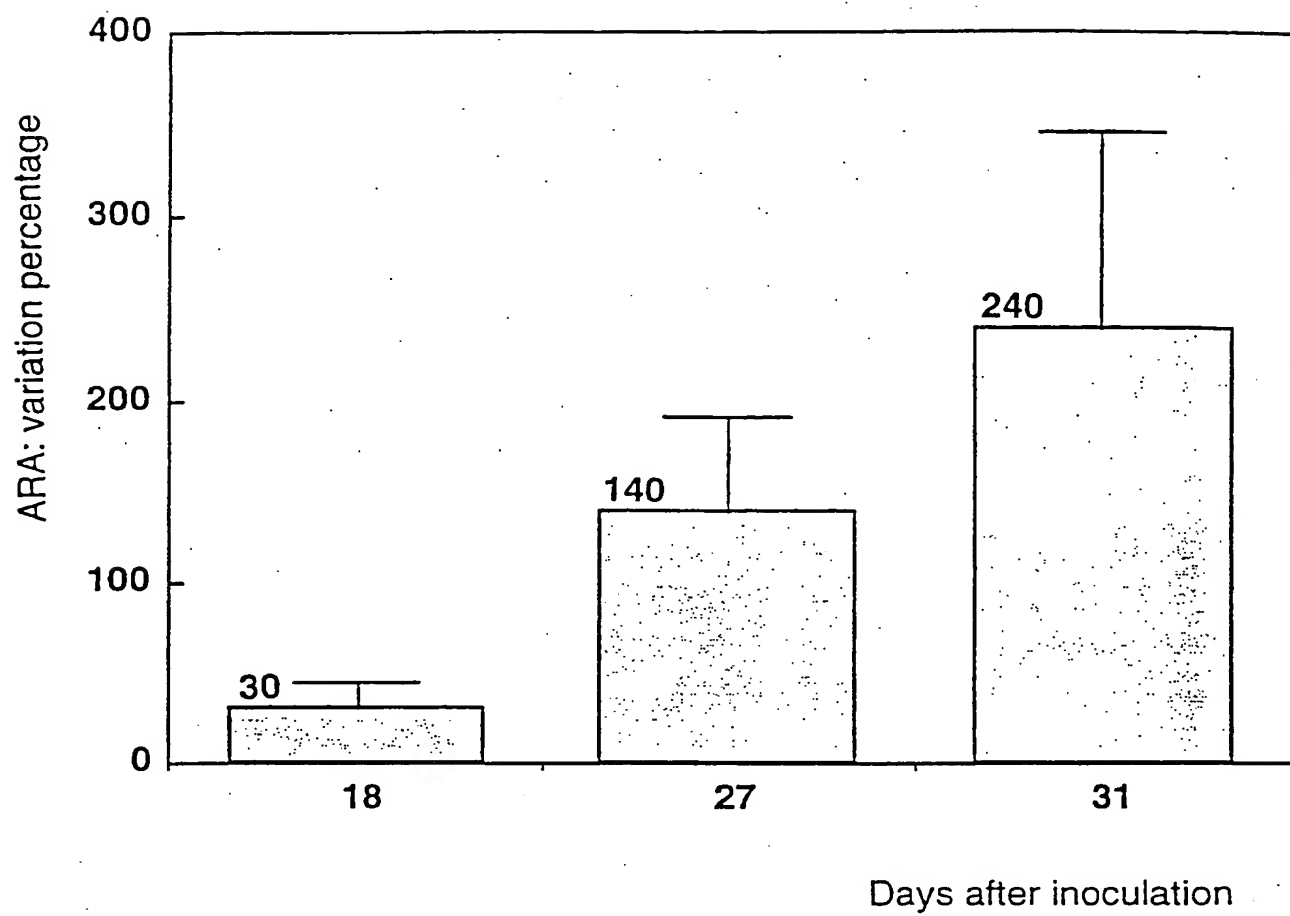


FIG. 6

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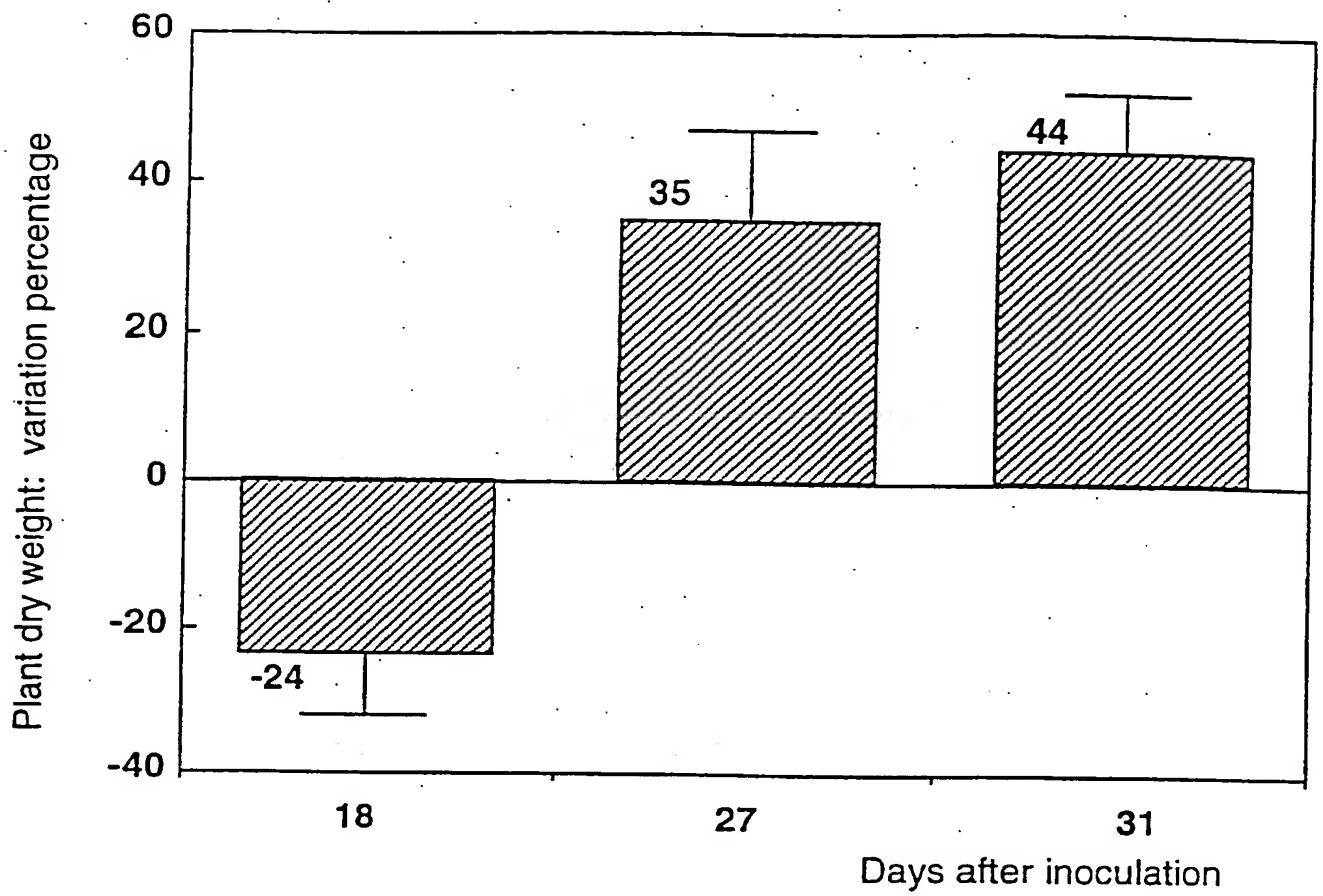


FIG. 7

GTGAGTGTGGTTGAGGTTCAATTATTACTATTTTGAAGCTGTGTATT-TCCCTTTTCTAATATGCACCTATTCAIGTTTCAGAAIG// rolA A4 coding region
s.s.

+1

└→

s.s.

σ^{70} CONSENSUS
SEQUENCE

TTGAAT____17bp____TATAAT
-35 -10

σ^{38} CONSENSUS
SEQUENCE

CTATACT

-10

CGGCAAGT

GEAR BOX
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FIG. 8

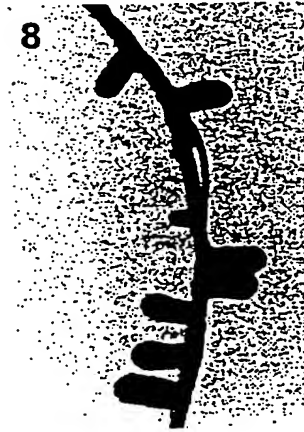
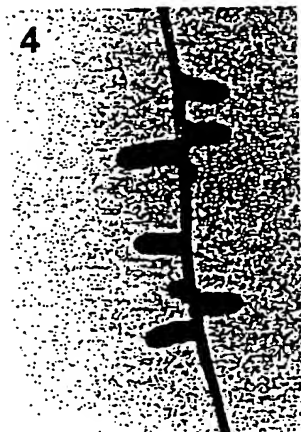
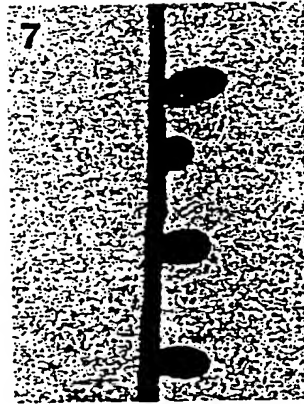
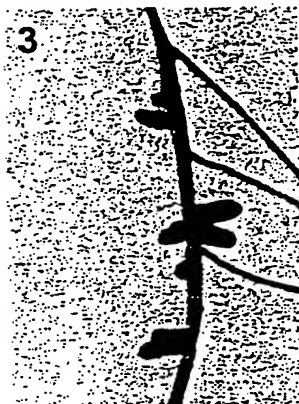
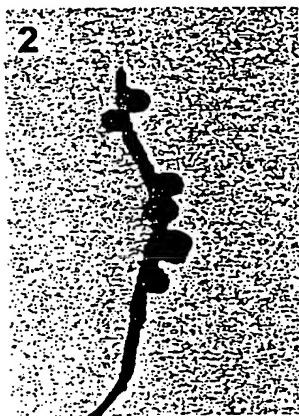
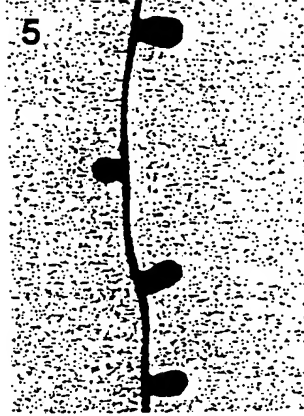
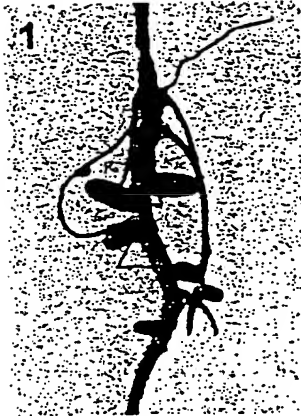


FIG. 9

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Days A. I.	Mnod(mg)/pp	
	RPR 1105	<i>iaaMtms2</i>
14	4 ± 1	5,0 ± 0,4
21	5 ± 1	7 ± 2
28	8 ± 2	12 ± 3
35	11 ± 1	12,2 ± 0,3
42	11 ± 4	23 ± 6
49	13 ± 3	18 ± 4

pendence	RPR 1105	0,26700	±	3,E-05
intercept	RPR 1105	0,26	±	0,03

pendence	<i>iaaMtms2</i>	0,4620	±	1,E-04
intercept	<i>iaaMtms2</i>	-1,69	±	0,1

FIG. 10A

mg of nodule/plant

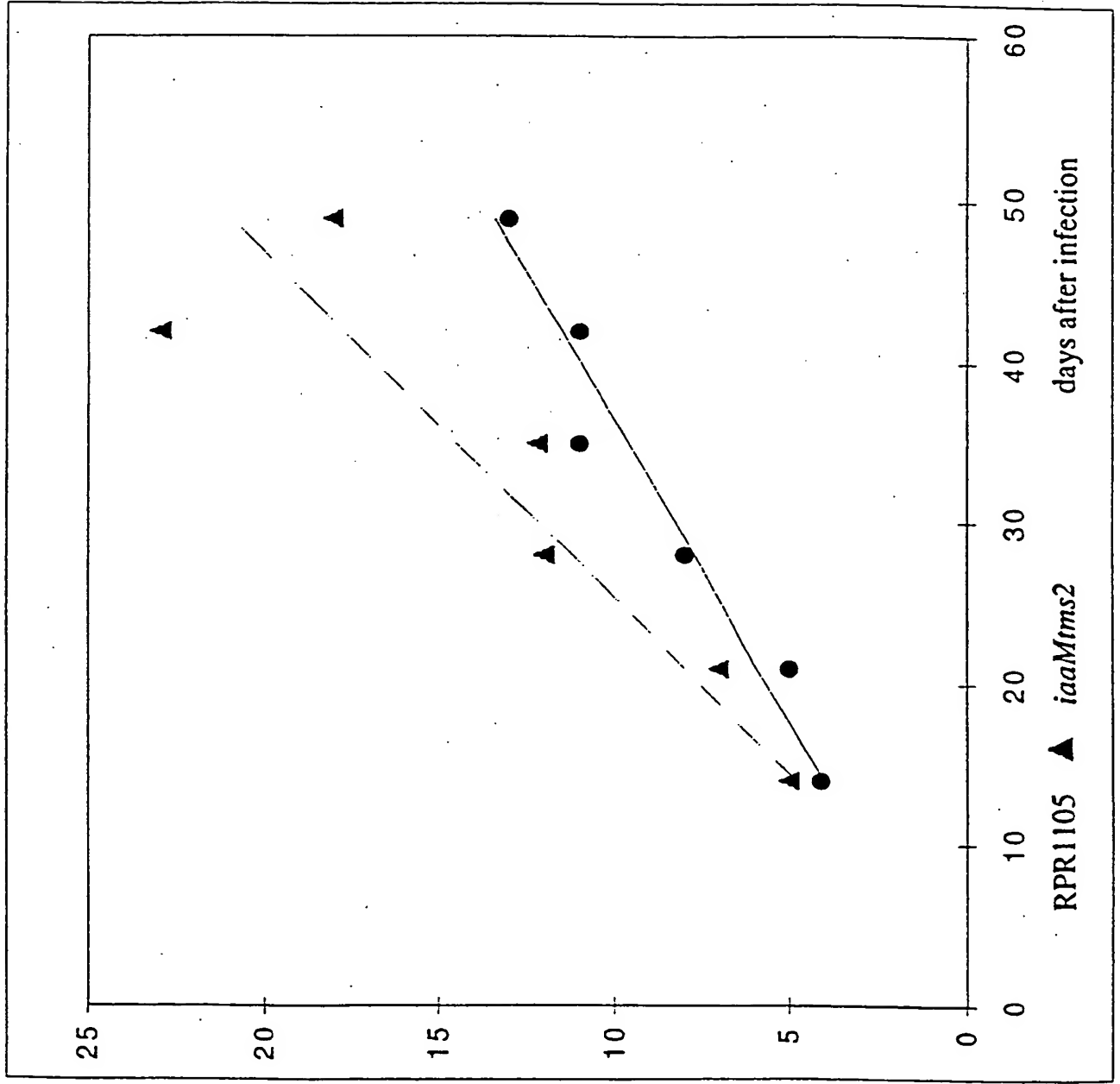
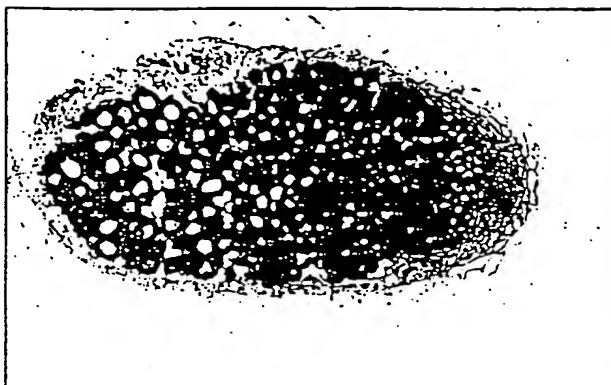
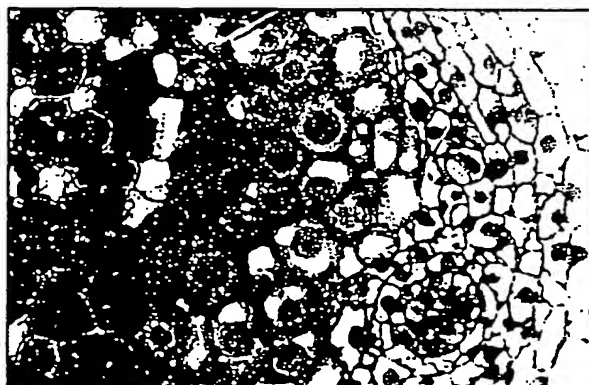


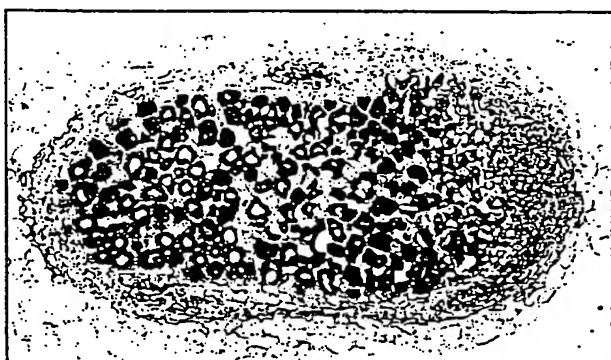
FIG. 10B



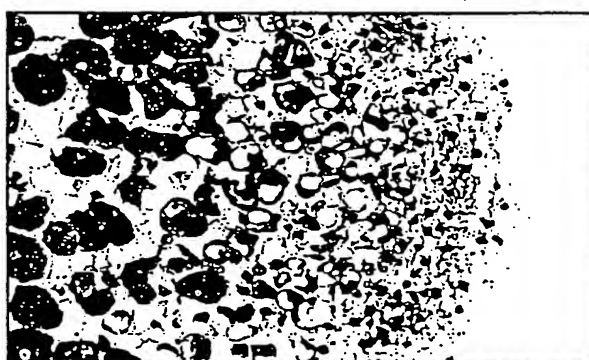
1



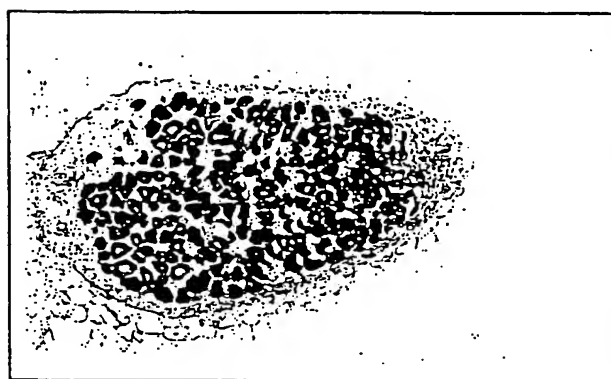
2



3



4



5

FIG. 11